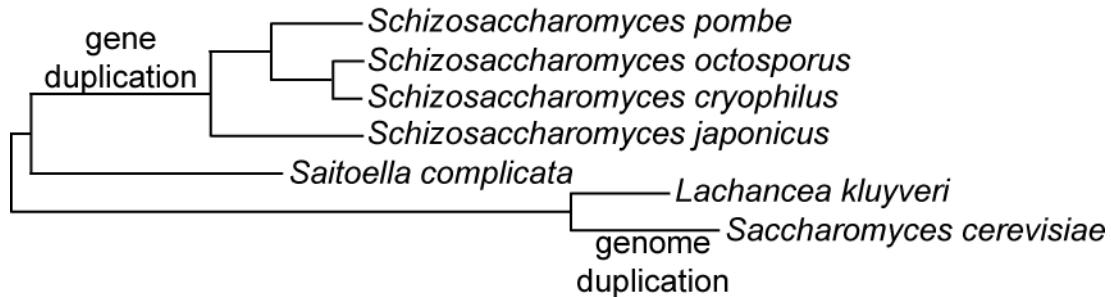


A.**B.**

Sequence alignment of Hbs1 and Sk1 proteins from various yeast species. The alignment shows conserved regions and two specific motifs, S1 and S2, highlighted in red and blue respectively. The alignment includes sequence identifiers, percentage identity, and the motif locations.

Protein Identifier	Identity (%)	Sequence	
pomb_Hbs1	1	MSRHRDVKNLDI DDYELDEEPG----EEELTEEQ E FRSAVATVRETLLGVP- I SEKEIA D TWYYYYFDVEKS V NYLQKASSKAGAKEKQNTDSQEK K Q-----	
japo_Hbs1	1	MSRH DKLNLD I NDYEDEFEP----EEELTEQ E AYRAE I EQVQTAEGTN- I PFQEIKDTWVYYFDVNKS V NYLLSKA-I K AK E KDAK K TPVAQSQT K -----	
octo_Hbs1	1	MSRHRNVKNL E I LEEYDA E P-----VEELT D EQ E QF R E A IAAV Q ET V EGLP-VTM K EIA D TWYYYYFDVEKS V NYLV K SCTARAE E EKG D ESHK K PP K -----	
cryo_Hbs1	1	MSRHRNIKNL D I AEYDY E DEPA---QE E LT D EQ E Q F R E A IAAV Q ET L D G LP-VTT K EIA D TWYYYYFDVEKS V NYLL K SCT T KAAA Q KE E SEKK K QN K -----	
scom_long	1	MSRHRDVRNLD E M YDDDDYYDG D G H DMTY E EQ E Q M E A GV V AAV H DAL N GVPGIT L KE I RE T LYYYYFD L E K SIAW L LE Q HSV V KPA A PK A PT A SK A -----	
pomb_Ski7	1	-----	
japo_Ski7	1	-----	
octo_Ski7	1	-----	
cryo_Ski7	1	-----	
scom_short	1	MSRHRDVRNLD E M YDDDDYYDG D G H DMTY E EQ E Q M E A GV V AAV H DAL N GVPGIT L KE I RE T LYYYYFD L E K SIAW L LE Q HSV V KPA A PK A PT A SK A -----	
pomb_Hbs1	98	-----	
japo_Hbs1	96	-----	
octo_Hbs1	96	-----	
cryo_Hbs1	98	-----	
scom_long	121	ISASVIIARHGITWGMGSTASIEPVHPPRVPPGGWGLGGSGK M SKLA A LAKARKEAQ-----	
pomb_Ski7	1	----- M SRSQLNS K KAK Q K--PPSEHPIGLSSIL K QDSSSSD S PNFFPSSTNDHQ E RT I NDTNFVVPE K Q K TS K LA L -----	
japo_Ski7	1	----- M SKLAK L SS K K A Q E NN T QQ N EE S KN V INS-----LTKS L ET S SI G SS S NT Q A T Q T R A P A V P SK L MS K LS Q -----	
octo_Ski7	1	----- M SKLSQLLN A K R T N Q N SS L SE S SS G K L TS L LK--SPSS S SE Q E H T A S V D T V H T T S N AA E SS A ST P Q P NG L SK L AA-----	
cryo_Ski7	1	----- M SKLSQLLN A K R AN Q NT P AP E SS G K L TS L LK--TPPPS V E-EHP A P L DF S Q T S N DT T E Y S A SS S Q P SG L SK L AA-----	
		motif S1	motif S1'
scom_short	101	-----	
pomb_Hbs1	98	-----	
japo_Hbs1	96	-----	
octo_Hbs1	96	-----	
cryo_Hbs1	98	-----	
scom_long	179	AASKAEKG A P G LA S V S LL S KL I AA K T A K P AAA P D T P E PPA-P Q A P A E P A V S E P E F S Q S S I P P S P S PP V ADD V DM S EA P LL V S P T P S S PE H P T F E PA L VE A T PS V F AT S L F -----	
pomb_Ski7	76	LAAERKKL H SSFP T QQ P P K T E K E K P I Q A H KK N V E ND F LL Q R F R K V R I A E K K -----SEQPSS H E I HL T DD D K T TL Q K Q M V --SDQL K NP Q E V K L A P P S S F A K C L T -----	
japo_Ski7	71	L Q K R QQ E QQ E K H AS L ASE T K N QS N SK V N D T A N L F S Q R K R L N SQ P TT P I I D S K Q SL N T E P V K A T P L S P I P H -TP S P PD R H N S C S T C Y --DH S I QT P D A S M Y M S F S F A K C L C -----	
octo_Ski7	75	LAAQKK N ENT E E T A N K R SL P SSSS--L Q T D S N K D S Y D K E Q L Q R F R K V R I A E K S P E N K P S L V H R Q E Q Q T P T R D T K L N I N ED D F A S P Q T S--SAHL F SS S P T S L M A P S V F A R C L -----	
cryo_Ski7	74	LAAQKK-NQNSEEASNKRPLPSSSS-LQPD T K S N D S F D K E Q L Q R F R K V R I A E K T P S E L D H D T L S R Y P A P P E Q V NN I Y D --PSS S E L P--QYD L SSSSP T S M M A P S I F A R C L -----	
scom_short	101	-----	

pomb_Hbs1 98 -----NKSKEALADAKDPLDESSNGIKNLISLNKDEPAFQTNGEVKMKNSSES DNQPEKKI IKKQNPTDLVS VPEIFEQSNPKPV
 japo_Hbs1 96 -----ENSTKDVTAAVKVTISQPSNALRKQVPIQTTKTESEHAQIPQKEPKL LDIPSLYEKEHEKPS
 octo_Hbs1 96 -----QEQQKQQTVEAQPSTKNASASLNVSMKKLNLNGKEQTGEKSDEETQTPRKQESSLLIDVPKIEESQPKPV
 cryo_Hbs1 98 -----QEQNSTAGNRPSSKKTQVLDASLEKLNLGKEQTQHETAKGLEETQVKVRQESSALIDVPKIEESHPKPV
 scom_long 298 GPEDYEVKYRSKRRREADSRFFYL PASAAPEVAKVFSGPSPDDVVIAAREEGPKSGKKAKTAKKDQAIAPTNQLEVDMQAMNMGSATGGAPAPAPPIKIARKKVNVEEYAKVDVKES
 pomb_Ski7 184 GAKKRVFEDQ-----IEIHLSSLLGFNA PSPDDIVLMAQS---KSKFQKHKRDLDEQLLNSVKS QQLPKQKNTNDSNDHTLILSQDQ---LIELSKLV----KPR
 japo_Ski7 188 FDIHKPPKDN-----IVIPYSRSALLGFHA PSPDDIVLAAQS---KSKAFQKTANK-----AAVGAAQPKPKEDVKLS---IEEITRHA----KPV
 octo_Ski7 190 GAKKKALQNE-----IKVNLKRSSILGFNT PSPDDIVLMAQS---KSKGFGQKHSKRDKA LHD SLEQFKAANLQQA SNSTSPISTKDV LIDKEK---LLEISKEV----KPT
 cryo_Ski7 188 GAKKKAMQKE-----IKVNLKRSSILGFNA PSPDDIVLMAQS---KSKGFGQKYNRKDRALFNSLQQLKNA SLOQTSTSIA TPTVTKDVMVDKSK---LLELSKDI----KPT

motif S3

scom_short 101 -----SAAPEVAKVFSGPSPDDVVIAAREEGPKSGKKAKTAKKDQAIAPTNQLEVDMQAMNMGSATGGAPAPAPPIKIARKKVNVEEYAKVDVKES

pomb_Hbs1 178 VHLVVTGHVDGKSTMLGRIMFELGEINSRSQMQLHNEAANSKGFSFYAWLLDTTEERARGVTMDVASTTFESDKKIYEIGDAPGHRFISGMIAGASSADFAVLVVDSSQNNFERGF
 japo_Hbs1 161 IHLVVS GHVDGKSTMIGRLLYEVGMVDERSMQKLQSAANA GKGFSFYAWLLDSTD EERARGVTMDVADTTFESSKHIYQIGDAPGHKFISGMIAGAYLSDYAILVV DASPNNFERGF
 octo_Hbs1 169 VHLVVTGHVDGKSTMLGRIMFELGDVNTRSMQKLHNEAANSQKGFSFYAWLLDSTD EERLRGVMTMDVAATS FESKKQIYEVG DAPGHKFISGMIAGAA SAEFAILVV DSSQNNYERGF
 cryo_Hbs1 169 VHLVVTGHVDGKSTMLGRIMYQLGDVNSRSQMQLHNEAANQKGFSFYAWLLDSTD EERLRGVMTMDVAATS FESKKQIYEVG DAPGHKFISGMIAGAA SAEFAILVV DSSQNNYERGF
 scom_long 418 ANFVII GHVDAGKSTMMGRLLY DIGAVDERTIQKFRKESEKM GKGFSF ALAWVMDSTD EERARGVTVDIATNQFETPKRKFTILDAPGHADFVPNMIAGAAQADFAVLVIDASTGGFESGF
 pomb_Ski7 281 TKLILLGPPKSGKTKTLLSRLFFQIGSFDPKTMQKCTVLNAK---KE SLSV LKSTKTKWYDFETFSNSYSSTI IDFPLGI FTTNASS--RDNF LKHSSLFQVMNTAIFTIDCLNP----L
 japo_Ski7 264 IHIGIFGDVGAGKKSLLSRLFLYQIGGLDTKHAQKCRLLNSR---QCTMESI LSQSN DWNFETFASCCTLLS YSEL-----DINACSRRLFLD MGVFVLRPHPTD----
 octo_Ski7 287 TRLSIFGPPKV GKTLLARLLYQVGALDI KMQRCALLNSR---KENFSSV LDRDAS GLYQFETFSHNYLSSLFALPL-----QD LASFAPFLQT DIVI VII HAKYP----L
 cryo_Ski7 285 NRISIFGAPKV GKTLLARLLYQVGALDI KMQRC TLLNSR---KENLSSV LGRNES GLYQFETFSHNYLSSLFALPL-----HD LAA LATFLQTTDIAVIVI HAKYP----L

motif G1

motif G2

motif G3

scom_short 195 ANFVII GHVDAGKSTMMGRLLY DIGAVDERTIQKFRKESEKM GKGFSF ALAWVMDSTD EERARGVTVDIATNQFETPKRKFTILDAPGHADFVPNMIAGAAQADFAVLVIDASTGGFESGF

pomb_Hbs1 298 LENGQTREHAYLLRALGISEIVVSVNKLDLMSWSEDRFQEIKNIVSDFLIKMVGFKTSNVHFVPI SAISGTNLIQKDSSDLYKWKYKGPTLISALDQLVPP---EKPYRKPLRLSIDDVYR
 japo_Hbs1 281 FSNGQTREHAYLLRALSVKGIAVCVNKLDTVWSYERFIAIKENI LD FLVSKVGF KETMVHIVPVSGLSEN LIK RDEPKL LSWYNGPTLMNI LDDFVPP---TKPVKASLRITVNDTYR
 octo_Hbs1 289 LANGQTREHAYLLRALGVSELA VAVVNKL DLSWSSLD RYNEIKASVSDFLIRMVGFKEENVHFIPVSAVGVN LIKQS PPLYTWYNGPTLIEVLDNFNPP---LKAYRGPLRV SVHDTYR
 cryo_Hbs1 289 LANGQTREHAYLLRALGVSEVA VAVVNKL DLSWSSLD RYFEIKSISDFLVRMVGFKEENVHFIPVSAVGVN LIKQS PPLYTWYNGPTLIEVLDNFNPP---LKSYRGPLRV SVHDTYR
 scom_long 538 NVRGQTKEHALLVRS LGVQNLIVAVNKLDSDVN HRFEEIEMQVSQFLTN-AGFD PQNVQYI PC SGLTGENLVKRS AEPALTWFNGPTV LGALESIA PT---ARAIEKSLRISVQDVYK
 pomb_Ski7 392 EGLDGII SILQIMNGLISSYMF AITK MDEIEW D ENKFINLVNSI QSF LKESC GIEK-SK FIPISGLKG TNLTSI S QEK L SQWY KSDT LLG KIDKEADTN GTWNFL NLPLS LTISHI
 japo_Ski7 369 TELQHICEVVRLDVLQI RLFVFI ITGMDI VDWSE SAFT ETCOLL QQAIQKQCKQ SI PATQFVPT S ALNGENL TSLSQG KLR KWY RSGT VLSV L DELCESI LAPQ K KALNKEPLRASLESI
 octo_Ski7 393 DEIESIVYLLRLCYGEKIRDVLFVV TQMDQV SWQEEQYQFAVSS ITCLKDTHG ITVSPSS FIPVSGF KGDNV TTS FGT LQ SWY GS DTLLG KIDEL SDN STDE KIQLQH LPLS LSIT SW
 cryo_Ski7 391 DEIESIVC LRLF YGVTI K N VLFVV TQMDQI SWEEEQYQYV VSSVIT CLK ETY SITV PWSS FLPV SGF FRGD NL TTLS YGT LQ SWY DDE TL L GKID EL SDK S IDE K L QLQH LPLS TIT SW

motif G4

motif G5

scom_short 315 NVRGQTKEHALLVRS LGVQNLIVAVNKLDSDVN HRFEEIEMQVSQFLTN-AGFD PQNVQYI PC SGLTGENLVKRS AEPALTWFNGPTV LGALESIA PT---ARAIEKSLRISVQDVYK

pomb_Hbs1 416 S-----PRSVTVTGRVEAGNVQVNQVLYDVSSQEDAYVKN V IRNSDPSST WAGDTVTLQLADIEVNQLRPGDILSNYENPVRRV SFVAEIQTFDIHGFILSGSTLVLH GR T--V
 japo_Hbs1 398 S-----AKGVCVQGRV ESGNVQNNQVLFNV STNSDAYVKSIIRNGHP-HDWSVAGDNITMQLTDIDANEIRPGDILTTTSQPVKKT KSFIAEIQTFDILRPI LSGATLVLH RGRL--S
 octo_Hbs1 406 S-----VRGATISGRV ESGN I QNGQVLYNVSSQEDAYVKAVMRNSDPSV SWAVAGDNVTVQLT DIEPNQISLGDILSTYSDPV KRS KSF TADI IQTFDILG PI LSGSTLVLH GR T--V
 cryo_Hbs1 406 S-----VRGATISGRV ESGN I QNGQVLYNVSSQEDAYVKAVMRNSDPSV SWAVAGDNVTVQLT DIEPNQISLGDILSTYNDPV KRT KSFIAEIQTFDILG PI LSGSTLVLH GR T--V
 scom_long 654 AGVT----GGSVTISGRV DAGNVQVGETV HAAPS GEPA TVKS -M QVNDIAD WAGSNV VLNLDIDPMHL KAGDILCDPLNPVPTV RAF RARI ITF DLAR PITNGATIVLH GR RINEA
 pomb_Ski7 511 TPLP----ENQSHIYCSIHSGMLQDSQKLYVGTGRLETQITG-LSSDENPKGFVNAGDMIQAKIPTLP--NLC PGI LADSIDAFTSSK TAYNATWF HGSLEKGKSMH VIALFG CHAVL
 japo_Ski7 484 QLEKTGESIGEVNATALI HSGI LQDMPF YLGAKLACTVTT THSITN SHAFGV SGERVRLSFTLSSN-CPPVGSF ISG DYAVTHSS RVLY FQGIA LQP WMS-MDV HTMDL FVCGYHCK
 octo_Ski7 508 SLLP----ENKIRAEFY VHSGI LQAHQS I YTSVGK I ETKV HG-I KLRQH ERTW C LPGE QTEM HLT SP--NLNGTLCVDEENSY HLSRNAYITVFSIDSSLKAQNPVY VNAFF GA F R MS
 cryo_Ski7 506 SLLP----ENKIAAQFDV HSGIVQAYQK I YTSVGK LEARV QS-LK LHQHARTW C LPGE HTEM HLT SP--NLNGTLCIDVENSY HLSRIAYISAFTFDSSLKVH SPI VNAFF GA F S MN

scom_short 431 AGVT----GGSVTISGRV DAGNVQVGETV HAAPS GEPA TVKS -M QVNDIAD WAGSNV VLNLDIDPMHL KAGDILCDPLNPVPTV RAF RARI ITF DLAR PITNGATIVLH GR RINEA

pomb_Hbs1 528 TSVSLKIVTVN-----NKRSRHIA SRKRALV RISFLDGLFPLCLAECPALGRFILRRS--GDTVAAGIVKELC
 japo_Hbs1 508 APVVIK L QSVN-----GKKV RHITS RKT AIVKFTFLDGSYPICTVDDCKT L GK II L RRE--GDTVAAGIVRKV--
 octo_Hbs1 519 APVAVKII VN-----GKKR HITS RKR ASM QVN FLEG LYPV CTSEESP SLSR I I L RRD--GNTIAAGIVK T IL-
 cryo_Hbs1 519 ASVSVKII SVN-----GKKR HITS RKR ASI QVN FLDGSYPVCPSEESS SLSR I I L RRD--GNTIAAGIVK T IL-
 scom_long 769 ARIQALVATIDRADGQI I KKKPRHLASGQS AVVEIAFLGNGI PMETFKDSKDLGRV I RTG--GDTIAAGIVDEL F-
 pomb_Ski7 624 TKLYCFTDSQ EKA---PNAIGNDLE RNR TSLV KIELEN AFPLV KES YINTL SRLV F SEKW NSLIAFGTV LSLHD
 japo_Ski7 602 ARLSIMSTERE-----STQ QEDGP LEFL KAEI LDQ PIVACTS STLL SLSR I I AGNV DDH TVN VI GF IT SML-
 octo_Ski7 621 GKVFLYSEGKEND----M ALR GIEFH SKD LLLR LE LDQ S VP LVEHN NVP SLSR I I LLSKEE N ALLA SGV VL SV QK
 cryo_Ski7 619 AKVFFYSEGKENT----M ALQGLQF QN KNLFLIR L LDQ PIVPL VEFN T VP SLSR V I LLSKEAKT L WA SGII ISV QK

scom_short 546 ARIQALVATIDRADGQI I KKKPRHLASGQS AVVEIAFLGNGI PMETFKDSKDLGRV I RTG--GDTIAAGIVDEL-